

PCT10

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PATENT APPLICATION: US/10/009,340

DATE: 07/16/2002

TIME: 14:02:31

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Output Set: N:\CRF3\07162002\J009340.raw

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4 <110> APPLICANT: DUBREUCQ, BERTRAND
         LEPINIEC, LOIC
5
         CABOCHE, MICHEL
6
8 <120> TITLE OF INVENTION: PROMOTER ENABLING TRANSGENE EXPRESSION IN THE WHOLE
        PLANT EXCEPT IN THE SEED
11 <130> FILE REFERENCE: 065691-0262
13 <140> CURRENT APPLICATION NUMBER: 10/009,340
14 <141> CURRENT FILING DATE: 2001-12-20
16 <150> PRIOR APPLICATION NUMBER: FR 99/07362
17 <151> PRIOR FILING DATE: 1999-06-10
19 <150> PRIOR APPLICATION NUMBER: PCT/FR00/01574
20 <151> PRIOR FILING DATE: 2000-06-08
22 <160> NUMBER OF SEQ ID NOS: 5
24 <170> SOFTWARE: PatentIn Ver. 2.1
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 932
28 <212> TYPE: DNA
29 <213> ORGANISM: Arabidopsis thaliana
31 <220> FEATURE:
32 <223> OTHER INFORMATION: FAH promoter in Arabidopsis thaliana.
34 <400> SEQUENCE: 1
35 cagctgtagc atcttgatat tgctgatact cagccacaag atcgttcatg ttactctctg 60
36 cttcattaaa ctccatctcg tccattcctt cttctgtgta ccaatgcaag aaagcttatc 120
37 tcaacatcag gctgatataa ccaatatctt acttctttta catttgtgaa atggaaccaa 180
38 cccatttttc tqqaaaaagt gctaaccaaa catttgatta accgtatcac tactactttc 240
39 atttctatct tctgtttcat tatgctgact atttaagctc cgttgtcaaa tctctaagtt 300
40 agacataaaa gacaaagact aatcaattgt catcacacca gcgtcgtcga gtgagctata 360
41 ttaatcgtgg attttaagca ttaaagaaac attctatagt actaaagcaa ataaaataat 420
42 tataatcaaa cactatgctt gacactggtc acgtgtactg gtagtgaatg attctacatc 480
43 ataagaggcc gcatcaaaat cctaaaaata agcataatga attaatcatt tacaaatttt 540
44 attttactca ataagaaaat cgaaagtatg attattatct agctgccaca atcttcgaat 600
45 ttaatattta ctcaagaaga gaccgacttt aatccttgac tttctcattg ctctatggaa 660
46 aatgattaaa gcagtcaata aaatcttttg acattgttgg cagaagacca ataattcgaa 720
47 gtctaaaatg taatcgtcca cacagtgtat gagtatccta gtatttttt tcttttccat 780
48 ataagttgaa tttgtaatat atatagtgta atgttgttta tttgtggcaa cgtacaaaat 840
49 tgggaatcct ataagtgcga cgacaagtga caagacgagg ctatgaacag ctaatgtatg 900
                                                                      932
50 aaqaqaqcca aaaqaqcaac aacctggcac ag
53 <210> SEQ ID NO: 2
54 <211> LENGTH: 20
55 <212> TYPE: DNA
56 <213> ORGANISM: Artificial Sequence
58 <220> FEATURE:
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59 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer

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62 ttcatgttac tctctgcttc
                                                                      20
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66 <211> LENGTH: 23
67 <212> TYPE: DNA
68 <213> ORGANISM: Artificial Sequence
70 <220> FEATURE:
71 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
73 <400> SEQUENCE: 3
74 qqaaaqqaaa caaatacqqa ttc
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77 <210> SEQ ID NO: 4
78 <211> LENGTH: 237
79 <212> TYPE: PRT
80 <213> ORGANISM: Arabidopsis thaliana
82 <220> FEATURE:
83 <223> OTHER INFORMATION: Fatty acid hydroxylase Fah 1p
85 <400> SEQUENCE: 4
86 Met Val Ala Gln Gly Phe Thr Val Asp Leu Lys Lys Pro Leu Val Phe
89 Gln Val Gly His Leu Gly Glu Asp Tyr Glu Glu Trp Val His Gln Pro
                                    25
92 Ile Ala Thr Lys Glu Gly Pro Arg Phe Phe Gln Ser Asp Phe Trp Glu
            35
                                40
95 Phe Leu Thr Leu Thr Val Trp Trp Ala Val Pro Val Ile Trp Leu Pro
                            55
98 Val Val Trp Cys Ile Ser Arg Ser Val Ser Met Gly Cys Ser Leu
                        70
101 Pro Glu Ile Val Pro Ile Val Val Met Gly Ile Phe Ile Trp Thr Phe
102
                     85
                                         90
104 Phe Glu Tyr Val Leu His Arg Phe Val Phe His Ile Lys Thr Lys Ser
                                    105
107 Tyr Trp Gly Asn Thr Ala His Tyr Leu Ile His Gly Cys His His Lys
108
           115
                                120
                                                    125
110 His Pro Met Asp His Leu Arg Leu Val Phe Pro Pro Thr Ala Thr Ala
                            135
                                                140
113 Ile Leu Cys Phe Pro Phe Trp Asn Ile Ala Lys Ala Ile Ser Thr Pro
114 145
                        150
                                            155
116 Ser Thr Ala Pro Ala Leu Phe Gly Gly Met Leu Gly Tyr Val Met
                    165
                                        170
119 Tyr Asp Val Thr His Tyr Tyr Leu His His Ala Gln Pro Thr Arg Pro
120
                                    185
122 Val Thr Lys Asn Leu Lys Lys Tyr His Leu Asn His His Phe Arg Ile
123
                                200
           195
125 Gln Asp Lys Gly Phe Gly Ile Thr Ser Ser Leu Trp Asp Ile Val Phe
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128 Gly Thr Leu Pro Thr Thr Lys Ala Pro Arg Lys Glu Gln
129 225
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132 <210> SEQ ID NO: 5
133 <211> LENGTH: 714
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134 <212> TYPE: DNA

135 <213> ORGANISM: Arabidopsis thaliana

137 <220> FEATURE:

138 <223> OTHER INFORMATION: Coding sequence for fatty acid hydroxylase

139 Fah 1P

141 <400> SEQUENCE: 5 142 atggttgctc agggattcac tgtggatctt aaaaagcccc ttgtatttca ggttggtcat 60 143 cttggagaag attatgagga atgggttcac caacctatcg cgaccaagga aggccctcgg 120 144 ttttttcaga gtgacttttg ggagttcttg acacttacag tttggtgggc agttcctgtc 180 145 atttggttgc cagttgtagt ctggtgcata tcaaggtcag taagtatggg atgttcactt 240 146 ccagaaatcg tcccaattgt tgtcatggga atattcatct ggacattttt tgaatacgtt 300 147 cttcaccggt tcgttttcca cataaaaacg aagagttact ggggaaacac tgcacactat 360 148 cttattcacg gatgccatca taagcacccg atggaccacc ttcggctcgt ctttcctcct 420 149 actgcaactg cgattttatg ctttccgttc tggaacattg cgaaggctat ctcaactcct 480 150 tcaaccgcac ctgcattgtt tggtggaggc atgctcggat atgtgatgta cgatgtcact 540 151 cattattacc ttcaccatgc ccaacctact agaccagtga ccaaaaatct caagaagtac 600 152 catttgaatc atcacttcag gattcaggac aaaggatttg gtataacttc gtcgttatgg 660

153 gacatagtet ttgggacaet teccaecaea aaageeecea gaaaagagea atag

VERIFICATION SUMMARY

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